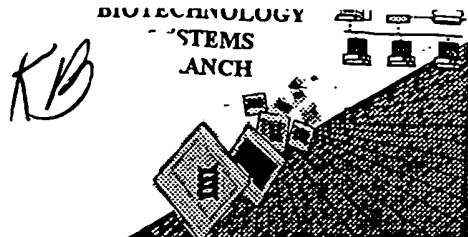PCT09
# CRF Problem Report

**The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):**

Application Serial Number: ___09|937,204A___

Filing Date: ___09/21/2001___

Date Processed by STIC: ___11/25/02___

## STIC Contact: **Mark Spencer, 703-308-4212**

### Nature of Problem:

The CRF (was):           Melted

___✓___ (circle one) (Damaged) or Unreadable (for Unreadable, see attached)

_____ Blank (no files on CRF) (see attached)

_____ Empty file (filename present, but no bytes in file) (see attached)

_____ Virus-infected. Virus name: _____ The STIC will not process the CRF.

_____ Not saved in ASCII text

_____ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.

_____ Did not contain a Sequence Listing. (see attached sample)

_____ Other:

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## PLEASE USE THE CHECKER VERSION 3.1 PROGRAM TO REDUCE ERRORS.
## SEE BELOW FOR ADDRESS:
### http:/www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)

2. **U.S. Postal Service:** U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202

3. **Hand Carry directly to:**
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

4. **Federal Express, United Parcel Service , or other delivery service to:** U.S. Patent and Trademark Office, B x Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

PCT09

RAW SEQUENCE LISTING                    DATE: 04/25/2002
PATENT APPLICATION:  US/09/937,204      TIME: 11:37:43

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

```
 4 <110> APPLICANT: Fritig, Bernard
 5         Toquin, Valerie
 6         Geoffroy, Pierrette
 7         Legrand, Michel
 8         Kauffmann, Serge
11 <120> TITLE OF INVENTION: INDUCIBLE COMTII PROMOTER, CHIMERA GENE
12         CONTAINING SAME AND TRANSFORMED PLANTS
15 <130> FILE REFERENCE: A34638-PCT-USA
17 <140> CURRENT APPLICATION NUMBER: US 09/937,204
18 <141> CURRENT FILING DATE: 2000-03-22
20 <150> PRIOR APPLICATION NUMBER: PCT/FR99/03700
21 <151> PRIOR FILING DATE: 1999-03-22
23 <150> PRIOR APPLICATION NUMBER: PCT/FR99/07646
24 <151> PRIOR FILING DATE: 1999-06-11
26 <160> NUMBER OF SEQ ID NOS: 26
28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
```

ERRORED SEQUENCES

```
712 <210> SEQ ID NO: 26
713 <211> LENGTH: 30
714 <212> TYPE: DNA
715 <213> ORGANISM: Artificial Sequence
717 <220> FEATURE:
718 <223> OTHER INFORMATION: Synthetic Oligonucleotide PS14
720 <400> SEQUENCE: 26
721 acgcgtcgac cagtggtgag tttagctgtc                          30
```
E--> 734 hy02:358522.1  -delete

VARIABLE LOCATION SUMMARY       DATE: 04/25/2002
PATENT APPLICATION:  US/09/937,204     TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

<u>Use of n's or Xaa's(NEW RULES):</u>
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
·residue n or Xaa represents.

Seq#:26; N Pos. 31

**VERIFICATION SUMMARY**　　　　　　　　DATE: 04/25/2002
PATENT APPLICATION:　US/09/937,204　　　TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

```
L:44 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:47 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:59 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:63 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:67 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:75 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:79 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:82 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:86 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:89 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:92 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:146 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:149 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:152 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:447 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
L:536 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:734 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:26
L:734 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:30
L:734 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:34 SEQ:26
L:734 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:734 M:112 C: (48) String data converted to lower case,
L:734 M:252 E: No. of Seq. differs, <211> LENGTH:Input:30 Found:34 SEQ:26
```